



## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09 1824,924A  
Source: 1600  
Date Processed by STIC: 12/10/02

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 3.1 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name,  
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,  
2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office,  
Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

RECEIVED

DEC 20 2002

TECH CENTER 1600/2900



1600

Does Not Comply  
Corrected Filing Date NeededRAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/824,924ADATE: 12/10/2002  
TIME: 13:38:43Input Set : A:\07039-235.TXT  
Output Set: N:\CRF4\12102002\I824924A.raw

4 <110> APPLICANT: Eckman, Christopher B.  
 5 Eckman, Elizabeth A.  
 8 <120> TITLE OF INVENTION: ENDOTHELIN CONVERTING ENZYMES AND THE  
 9 AMYLOID BETA PEPTIDE  
 11 <130> FILE REFERENCE: 07039-235001  
 13 <140> CURRENT APPLICATION NUMBER: 09/824,924A  
 14 <141> CURRENT FILING DATE: 2001-04-03  
 16 <150> PRIOR APPLICATION NUMBER: 60/233,012  
 17 <151> PRIOR FILING DATE: 2000-09-15  
 E--> 19 <160> NUMBER OF SEQ ID NOS: ⑦- found 8  
 21 <170> SOFTWARE: FastSEQ for Windows Version 4.0

## ERRORED SEQUENCES

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/824,924A

DATE: 12/10/2002

TIME: 13:38:44

Input Set : A:\07039-235.TXT

Output Set: N:\CRF4\12102002\I824924A.raw

L:109 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!  
L:113 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8  
L:117 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8  
L:121 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8  
L:126 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8  
L:131 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8  
L:132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0  
L:19 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (7) Counted (8)